

# Annotation Summary Results

[Help and Tool Manual](#)

Current Gene List: List\_1

Current Background: Homo sapiens

555 DAVID IDs

Check Defaults ☒

Clear All

- ☒ **Disease** (2 selected)
- ☒ **Functional\_Annotations** (6 selected)
- ☒ **Gene\_Ontology** (3 selected)
- ☒ **General\_Annotations** (0 selected)
- ☒ **Interactions** (1 selected)
- ☒ **Literature** (0 selected)
- ☒ **Pathways** (3 selected)

<input checked="" type="checkbox"/> <b>BBID</b>	1.1%	6	Chart	
<input checked="" type="checkbox"/> <b>BIOCARTA</b>	6.3%	35	Chart	
<input type="checkbox"/> EC_NUMBER	15.5%	86	Chart	
<input checked="" type="checkbox"/> <b>KEGG_PATHWAY</b>	33.5%	186	Chart	
<input type="checkbox"/> PATHBANK	6.5%	36	Chart	
<input type="checkbox"/> REACTOME_PATHWAY	46.8%	260	Chart	
<input type="checkbox"/> WIKIPATHWAYS	35.3%	196	Chart	

- ☒ **Protein\_Domains** (4 selected)
- ☒ **Tissue\_Expression** (0 selected)

\*\*\*Red annotation categories denote DAVID defined defaults\*\*\*

## Combined View for Selected Annotation

Functional Annotation Clustering

Functional Annotation Chart

Functional Annotation Table